

MACLEAN et al
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herpes simplex virus type 1 which has a non-functional $\gamma 34.5$ gene in the long repeat region (R_L).

25. (Amended) A method according to claim 20, wherein the mutant virus is a herpes simplex virus type 1 which has been modified by deletion, insertion, substitution or rearrangement within the Bam H1 \mathfrak{s} restriction fragment of the R_L terminal repeat.

30. (Amended) A method treating a cancer in a human [mammal], which method comprises the step of administering to the said human [mammal] an effective amount of a mutant herpes simplex virus type 1 wherein the mutant virus is a mutant strain 17 virus which has a non-functional $\gamma 34.5$ gene in the long repeat region (R_L).

36. (Amended) A method according to claim 30, wherein the mutant strain 17 virus has been modified by deletion, insertion, substitution or rearrangement within the Bam H1 \mathfrak{s} restriction fragment of the R_L terminal repeat.

IN THE ABSTRACT

Please add the abstract attached.

REMARKS

Reconsideration of this application is requested. Claims 20, 22-30 and 32-42 are in the application. Claims 21 and 31 have been cancelled while claims 20, 25, 30 and 36 are amended; the significance of these changes is discussed below. Claims 23 and 35 stand withdrawn from consideration.